# Hockey Pool Linear Regression

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### **Data Preparation and Library Import**

In this section, we begin by setting up our environment and preparing the data for analysis. The steps involved are as follows:

Installing and Loading Necessary Libraries

We start by installing and then loading the car package. This package provides functions for regression diagnostics and other statistical tools which are essential for our analysis.

### Loading the Dataset

Next, we load our datasets from CSV files. Two datasets are used in this analysis:

QuantHockey Dataset - Less Features LR.csv: This dataset contains selected features and is used for the QuantHockey Dataset Test.csv: This dataset is used for our predications.

```
class(df)

## [1] "function"

df <- read.csv("QuantHockey Dataset - Less Features LR.csv")

test_df <- read.csv("QuantHockey Dataset Test.csv")</pre>
```

## Regression Model Building

The steps include:

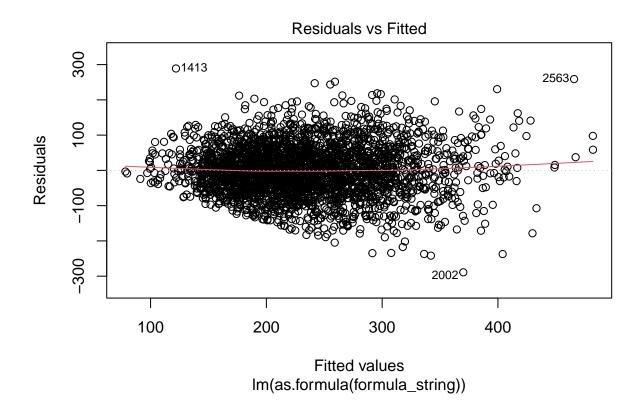
Renaming columns, defining the response variable, and fitting the model.

- Selection of Regressors: A subset of variables is chosen as regressors based on their potential influence on the response variable.
- Fitting the Linear Regression Model: Fit the linear regression model to our data.
- Model Summary and ANOVA: The summary of the regression model provides insights into the fit and ANOVA is performed to compare models or test certain model assumptions.

```
colnames(df) <- c("Rk", "Age", "GP", "G", "A", "P", "PlusMinus", "PPG", "SHG", "GWG", "OTG",
                 "PPA", "SHA", "GWA", "PPP", "SHP", "GWP", "OTP", "PPPPercent", "GoalsPer60",
                 "AssistsPer60", "PointsPer60", "ESGoalsPer60", "ESAssistsPer60", "ESPointsPer60",
                 "PPGoalsPer60", "PPAssistsPer60", "PPPointsPer60", "GoalsPerGame", "AssistsPerGame",
                 "PointsPerGame", "Shots", "ShootingPercent", "Hits", "BS", "Year", "FantasyPoints",
                 "NextYearFantasyPoints")
# Define the response variable
y <- df$NextYearFantasyPoints
# Fit a linear regression model with selected regressors
selected_regressors <- c("Rk", "Age", "GP", "G", "A", "PPG",</pre>
                       "PPA", "SHA", "GWG",
                       "Shots", "Hits", "BS")
# Include only the selected regressors in the formula
formula_string <- paste("y ~", paste(selected_regressors, collapse = " + "))</pre>
# Fit a linear regression model with selected regressors
model <- lm(as.formula(formula_string), data = df)</pre>
# View regression summary
summary(model)
##
## Call:
## lm(formula = as.formula(formula string), data = df)
## Residuals:
##
      Min
               1Q Median
                               30
                                      Max
## -289.07 -47.72 -2.18
                            46.26 288.99
##
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 335.09451 13.56131 24.710 < 2e-16 ***
                          0.03256 -5.301 1.24e-07 ***
## Rk
               -0.17258
               -3.06351
                           0.32213 -9.510 < 2e-16 ***
## Age
                          0.14872 -16.561 < 2e-16 ***
## GP
               -2.46297
## G
                0.88713
                          0.37929 2.339 0.019408 *
## A
                          0.27232 3.493 0.000486 ***
                0.95110
                          0.61378 2.533 0.011354 *
## PPG
                1.55487
## PPA
                ## SHA
                5.00185 1.93301 2.588 0.009714 **
                          0.90766 1.663 0.096488 .
## GWG
                1.50914
## Shots
                0.60340
                          0.04363 13.830 < 2e-16 ***
## Hits
                0.20134
                           0.03032 6.639 3.76e-11 ***
## BS
                0.14194
                           0.04666 3.042 0.002370 **
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## Residual standard error: 71.56 on 2834 degrees of freedom
## Multiple R-squared: 0.4175, Adjusted R-squared: 0.4151
## F-statistic: 169.3 on 12 and 2834 DF, p-value: < 2.2e-16
```

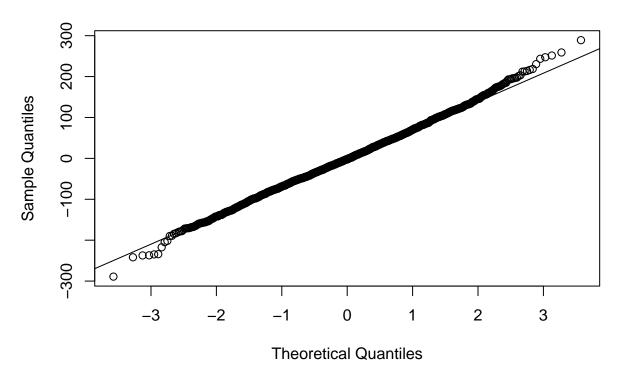
### anova(model)

```
## Analysis of Variance Table
## Response: y
             Df
                Sum Sq Mean Sq F value
             1 7023910 7023910 1371.7706 < 2.2e-16 ***
## Rk
              1 488148 488148 95.3354 < 2.2e-16 ***
## Age
## GP
             1 51579 51579 10.0735 0.001520 **
## G
             1 434727 434727 84.9022 < 2.2e-16 ***
             1 490117 490117 95.7200 < 2.2e-16 ***
## A
             1 152181 152181 29.7210 5.418e-08 ***
## PPG
             1 123088 123088 24.0392 9.972e-07 ***
## PPA
## SHA
             1 69448 69448 13.5632 0.000235 ***
                 20664 20664 4.0356 0.044644 *
## GWG
             1
## Shots
             1 1232403 1232403 240.6884 < 2.2e-16 ***
## Hits
             1 268524 268524
                                52.4427 5.681e-13 ***
                                9.2550 0.002370 **
                  47389 47389
             1
## Residuals 2834 14510999
                          5120
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
residuals <- resid(model)</pre>
# Residual Plot:
# Residual plot to visually inspect the distribution of residuals and identify
plot(model, which = 1)
```

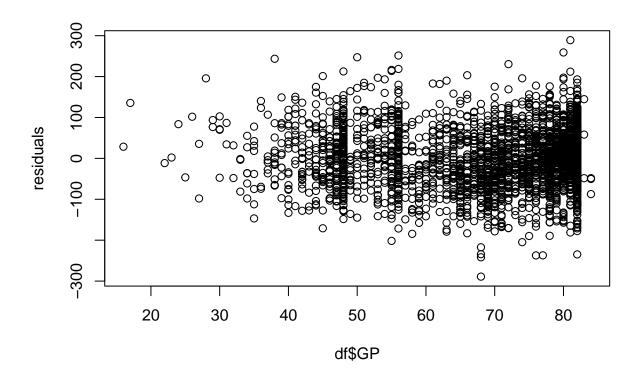


```
# Normality Check:
# Check for the normality of residuals using a Q-Q plot:
qqnorm(residuals)
qqline(residuals)
```

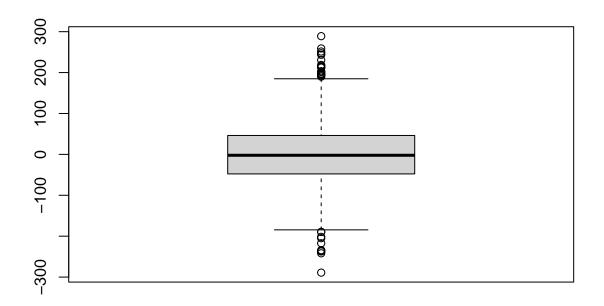
# Normal Q-Q Plot



# Homoscedasticity Check:
# Check for constant variance (homoscedasticity) by plotting residuals against the predictor variable
plot(df\$GP, residuals)

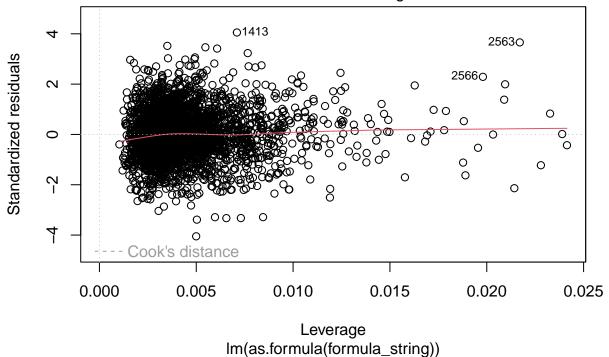


# Check for Outliers:
# Identify potential outliers by examining the residuals.
boxplot(residuals)



plot(model, which = 5) # Leverage plot

### Residuals vs Leverage



```
# Influence Points:
# Identify influential observations using Cook's distance:
infl <- influence.measures(model)
# plot(infl, which = 4) # Cook's distance plot</pre>
```

## Generating Predictions and Exporting Results

## **Making Predictions**

Using the predict() function, we generate predictions from our linear regression model for the test\_df dataset, which contains data we want to evaluate but wasn't used in the model training phase.

### Combining Predictions with Original Data

To facilitate easy analysis and comparison, we combine the predictions with the original test data. This merged dataset includes all the features from test\_df along with a new column Predictions that holds our model's predictions.

### Exporting the Results

Finally, we export the combined dataset with predictions to a CSV file. This file, named predictions.csv, can be used for detailed analysis, visualization, or presentation of the model's output.

```
predictions <- predict(model, newdata = test_df)

# Combine the original data with the predictions
result_data <- cbind(test_df, Predictions = predictions)

# Specify the file path for the CSV file
csv_file_path <- "predictions.csv"

# Export the combined data to a CSV file
write.csv(result_data, file = csv_file_path, row.names = FALSE)</pre>
```